

Selective depletion of fibroblasts preserves morphology and the functional integrity of peritoneum in transgenic mice with peritoneal fibrosing syndrome¹

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Selective depletion of fibroblasts preserves morphology and the functional integrity of peritoneum in transgenic mice with peritoneal fibrosing syndrome.

Background. A peritoneal fibrosing syndrome (PFS) can progressively reduce peritoneal ultrafiltration during chronic peritoneal dialysis in patients with renal failure. The pathogenesis of PFS is unclear and the role of peritoneal fibroblasts has not been evaluated experimentally.

Methods. We followed the fate of fibroblasts producing FSP1 in a mouse model using fibroblast-specific protein 1 (FSP1) as a marker. PFS was induced by daily peritoneal infusions of chlorhexidine gluconate (CHG) saline into transgenic mice expressing the thymidine kinase (Δtk) gene under the control of the *FSP1* promoter (FSP1. Δtk mice). To demonstrate the role of fibroblasts in PFS, we treated these FSP1. Δtk mice with a nucleoside analogue to induce DNA chain termination and fibroblast death.

Results. Mice receiving peritoneal infusions of CHG saline every other day for 2 weeks developed increasing numbers of FSP1⁺ fibroblasts in the subserosal layers of the visceral peritoneum. Mac-3⁺ monocytes (macrophages) subsequently accumulated over the next 2 weeks in association with increased deposition of type I collagen and increased endothelial vascularity (CD31⁺) in these subserosal tissues. Since these peritoneal fibroblasts expressed monocyte chemoattractant protein-1 (MCP-1), heat shock protein 47 (HSP47), and vascular endothelial growth factor (VEGF), we suspect they were partially responsible for macrophage recruitment, matrix production, and the neoangiogenesis in the subserosal tissue. Treatment of PFS in FSP1. Δtk transgenic mice with a nucleoside analogue selectively reduced the numbers of peritoneal fibroblasts and attenuated the attendant changes in peritoneal

histology. Rescuing the peritoneal membrane from chronic thickening and neoangiogenesis by reducing the number of fibroblasts also preserved ultrafiltration.

Conclusion. Peritoneal fibroblasts play a pivotal role in PFS, and their deletion using a fibroblasts-specific transgene was effective in preventing peritoneal fibrogenesis.

There is a tendency to lose peritoneal ultrafiltration and small molecule clearance [1–5] in patients receiving continuous ambulatory peritoneal dialysis (CAPD). Within a few months to a year after the initiation of CAPD, peritoneal membranes undergo a structural deterioration [2, 5–12] associated with membrane thickening and fibrosis [13]. This disorder is called the peritoneal fibrosing syndrome (PFS) [14]. Preserving the longevity of the peritoneal membrane and its relationship to normal peritoneal physiology is important to the long-term success of this therapy [1, 2, 5].

The expression of PFS is variable in humans, depending on individual susceptibility to membrane damage or the type of dialysate used for therapy. Long-term use of nonphysiologic dialysis fluids (low pH, hyperosmolarity, high glucose concentration, or high lactate), or contaminants derived from the nonenzymatic degradation of glucose, repeated bacterial peritonitis, or exposure to some drugs and antiseptics all contribute to PFS [2, 5, 14, 15]. The expected pathology in PFS seems independent of its cause [10, 12, 14, 16]. There have been several animal models of PFS, some of which are generated by substances currently unused for CAPD patients such as chlorhexidine [17–25]. However, these models share a number of similarities to human PFS [13] and have contributed to its understanding. The role of fibroblasts in the alteration of the peritoneum has never been studied in detail because of a dearth of fibroblast-specific markers.

Several years ago, we cloned fibroblast-specific protein 1

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(FSP1) [26, 27] and demonstrated the presence of fibroblasts in fibrosing tissues using anti-FSP1 antibodies [27–29]. FSP1⁺ fibroblasts remodel tissue architecture in organ fibrosis [27, 29]. When inflammation initiates tissue fibrogenesis, FSP1⁺ fibroblasts activate, proliferate, and produce fibrogenic molecules such as type I and III collagens, fibronectin, and proteoglycans [27, 29]. When tissue inflammation resolves, FSP1⁺ fibroblasts attenuate through apoptosis, leaving an acellular fibrous scar. With FSP1 as a marker, we studied a murine model of PFS to monitor the cell fate and pathophysiologic role of fibroblasts in the peritoneum challenged by chlorhexidine [30]. Transgenic mice (FSP1.Δtk) whose fibroblasts selectively express high levels of herpes simplex virus thymidine kinase (Δtk) under the control of the FSP1 promoter [26, 31] were treated with gancyclovir (GCV), a nucleoside analogue, which corrupts DNA synthesis by chain termination in dividing cells [31]. By generating PFS in FSP1.Δtk mice we asked if fibroblasts depletion would attenuate peritoneal histopathology or its functional deterioration.

We also focused on the role of macrophages in this study as they are a source of profibrotic molecules, including transforming growth factor-β1 (TGF-β1), platelet-derived growth factor (PDGF), tumor necrosis factor-α (TNF-α), interleukin-1 (IL-1), and fibronectin [29, 32]. These macrophages migrate to the peritoneal interstitium and peritoneal fluid from the circulation through small blood vessels [33]. The stimulus for this efflux is thought to be chemoattractant molecules, like monocyte chemoattractant protein-1 (MCP-1), regulated upon activation, normal T cell expressed and secreted (RANTES), macrophage inflammatory protein-1 (MIP-1), which are secreted by a variety of resident cells such as mesothelial cells, fibroblasts, or previously recruited macrophages [32, 34].

METHODS

Transgenic mice

FSP1.Δtk mice were housed individually with free access to chow and water and with humidity and temperature controlled under pathogen-free conditions. Animal care and treatment conformed to institutional guidelines. The creation of FSP1.Δtk mice and their breeding onto a Balb/c background has been previously described [31]. The FSP1 promoter directs the expression of Δtk exclusively in FSP1⁺ fibroblasts making these fibroblasts selectively susceptible to the lethal effects of nucleoside analogues either in culture or during experimental fibrosis [26, 31].

Generation of PFS in FSP1.Δtk or wild-type mice

Six 8-week-old male mice were maintained chronically with 21 gauge plastic catheters (Terumo, Tokyo, Japan) inserted from a position along the high back, through the

retroperitoneum, into the peritoneal cavity. Peritoneal fibrosis developed in peritoneal fibrosis mice following intraperitoneal injections of 0.03% chlorhexidine gluconate and 5% ethanol dissolved in 1.0 mL of saline (CHG saline) through a catheter every other day for 4 weeks. Control mice were treated with normal saline. Optimal conditions for producing chronic peritonitis with fibrogenesis were first determined in pilot studies (data not shown). In these early pilot studies, there were no substantial differences in pathology between the visceral and parietal peritoneum in peritoneal fibrosis mice with PFS at week 4. Since the surface area of visceral peritoneum is significantly larger and potentially more important as dialysis membrane, and since solid samples of bowel are easier to process, we used visceral peritoneum for our pathologic analyses. To minimize loss of mesothelium during tissue harvest, the rest of the intestines were immediately fixed in 4% paraformaldehyde [phosphate-buffered saline (PBS)] overnight, and then at least 10 small cylindric samples of bowel wall were randomly cut from fixed intestine and mounted in paraffin with the axis being kept vertical to the horizontal surface of the paraffin block. These tissue samples were used for histopathology and immunohistochemical analyses. The rest of the fixed bowel samples were rinsed in serial concentrations of sucrose solution and then snap-frozen for immunofluorescence.

Deletion of peritoneal fibroblasts in FSP1.Δtk or wild-type mice

To evaluate the effects of GCV on the deletion of fibroblasts in PFS, FSP1.Δtk (transgenic) or wild-type mice were divided into four comparative groups: peritoneal fibrosis-transgenic, control-transgenic, peritoneal fibrosis-wild-type, and control-wild-type mice. Peritoneal fibrosis-transgenic mice (*N* = 10) and peritoneal fibrosis-wild-type mice (*N* = 10) were treated with a catheter injection of 1.0 mL of CHG saline at 9:00 a.m. every other day for 4 weeks. Two weeks after initiation of CHG saline injections, GCV (kindly provided by F. Hoffmann, La Roche, Ltd., Basel, Switzerland) in PBS was also injected through the catheter (50mg/kg weight) at 9:00 p.m. every day for 2 weeks. Control-transgenic mice (*N* = 8) and control-wild-type mice (*N* = 9) received normal saline instead of CHG saline as above, and were likewise treated with GCV over the last 2 weeks of the experiment. During the study, mice losing a catheter were excluded from the data analysis (three mice from the control groups). Preliminary studies revealed that 2 weeks of GCV treatment did not yield any pathology in the peritoneal tissue of transgenic or wild-type mice (data not shown). All groups therefore received GCV in these experiments. At the end of the study, a modified peritoneal equilibrium test (PET) was carried out on each test subject and then tissue samples

were harvested for histopathologic evaluation and direct enzyme-linked immunosorbent assay (ELISA).

Flow cytometric analyses

Splenocytes were liberated from the spleens of peritoneal fibrosis-transgenic, peritoneal fibrosis-wild-type, control-transgenic, and control-wild-type mice by sieving through a nylon mesh. The cells were washed in PBS four times, and then incubated in hemolysis buffer (0.15 mol/L NH_4Cl , 1 mmol/L KHCO_3 , 0.1 mmol/L Na_2 ethylenediaminetetraacetic acid (EDTA) at pH 7.4) at room temperature for 5 minutes with occasional shaking [35]. The splenocytes were next washed in staining buffer [PBS with 1% bovine serum albumin (BSA)] four times and 50 μL aliquots of 10^6 cells were then incubated with various antibodies: monoclonal rat antimouse macrophages (Mac-3) (1:50) (PharMingen, San Diego, CA, USA), or fluorescein isothiocyanate (FITC)-conjugated rat antimouse CD4 or antimouse CD8 (PharMingen) for 20 minutes on ice; FITC-conjugated goat antirat immunoglobulin G (IgG) (1:100) was used for secondary staining of macrophage-labeled cells. All samples were then washed twice with staining buffer and fixed with 1% paraformaldehyde in PBS. The fixed cell samples were stored at 4°C under shade until scanned. FCAScan analysis (Becton-Dickinson, Franklin Lakes, NJ, USA) was performed on 10^4 cells using CellQuest software. Control cells were scanned with isotope-matched rat IgG and FITC-conjugated antibodies.

Histopathology, immunohistochemistry, and immunofluorescence

Four micron sections were cut from paraffin blocks and processed for hematoxylin-eosin staining or indirect immunoperoxidase staining. For the latter, after deparaffinization and rehydration, the sections were treated with proteinase K and boiled by microwave in citrate buffer for unmasking. Next, the sections were immersed in 3% H_2O_2 in methanol in order to inhibit endogenous peroxidase and nonspecific reaction products before flooding with 5% BSA in PBS. Rabbit anti-FSP1 (1:500) [27, 28], rabbit anti-type I collagen (1:400) [36], rat anti-Mac-3 (1:50), and rat anti-platelet-endothelial cell adhesion molecule-1 (PECAM-1) (CD31) (1:50) (PharMingen) were applied as primary antibodies followed by immunoenzyme reaction using biotin-conjugated antirabbit IgG or antrrat IgG (1:500) (American Qualex, San Clemente, CA, USA), and a Vectastain ABC Standard Kit (Vector Laboratories, Burlingame, CA, USA). Reaction development was accomplished using diaminobenzidine (DAB) as substrate (for detection of CD31, more intensive digestion and longer microwave exposure were needed). Ten visceral peritoneal fields at 150 \times magnification were assessed quantitatively in each immunostained section using a color image computer analyzer

(Mac SCOPE, version 2.5, Mitani Corp., Hukui, Japan). Immunohistochemical stainings were expressed as % area of reaction product within the subserosal stroma between the mesothelium and intestinal smooth muscle layer, including the mesothelium. Single or dual immunofluorescence was performed on 4 μm cryostat sections. The primary antibodies were employed as follows: goat anti-heat shock protein 47 (HSP47) (1:100) (Santa Cruz Biotechnology, Santa Cruz, CA, USA), anti-MCP-1 (1:100) (Santa Cruz Biotechnology), anti-vascular endothelial growth factor (VEGF) (1:100) (Santa Cruz Biotechnology), rabbit anti-FSP-1 (1:500), anti-tk protein (1:500) (generous gift from W.C. Summers, Yale University) conjugated with FITC using a FITC Protein Labeling Kit (Boehringer Mannheim GmbH, Mannheim, Germany), rat anti-Mac-3 (1:50), or FITC-conjugated rat antimouse CD4 and antimouse CD8 were used; a secondary antibody (Rhodamine-conjugated antigoat IgG and antirabbit IgG (1:500) (Chemicon International, Temecula, CA, USA) or FITC-conjugated anti-rabbit IgG and anti-rat IgG (1:500; American Qualex)) followed where necessary. These sections were analyzed under confocal microscope (MRC600) (Bio-Rad Laboratories, Hercules, CA). All of the secondary antibodies were isolated by immunoaffinity chromatography and absorbed for dual labeling. Control measures included omitting the primary antibody or substituting the primary antibody with species-specific normal IgG.

Detection of apoptotic cells

DNA fragmentation within apoptotic cell nuclei was detected using the MEBSTAIN Apoptosis Kit Direct (Medical and Biological Laboratories, Nagoya, Japan). After deparaffinization and treatment with proteinase K, histologic sections were incubated for 1 hour at 37°C with FITC-deoxyuridine triphosphate (dUTP) and terminal deoxynucleotidyl transferase, the latter linking FITC-dUTP to the 3'-OH fragmented ends of DNA. Immunocytochemistry with anti-FSP1 antibody and Rhodamine-conjugated antirabbit IgG was performed to colocalize apoptosis with FSP1 expression.

Indirect ELISA

Some of the small bowel tissues were sonicated in a cold radioimmunoprecipitation assay (RIPA) lysis buffer [1% nitroprusside-40, 0.1% sodium dodecyl sulfate (SDS), 100 $\mu\text{g}/\text{mL}$ phenylmethylsulfonylfluoride, 0.5% sodium deoxycholate, 1 mmol/L sodium orthovanadate, 2 $\mu\text{g}/\text{mL}$ aprotinin, 2 $\mu\text{g}/\text{mL}$ antipain, and 2 $\mu\text{g}/\text{mL}$ leupeptin in PBS), and the homogenates were centrifuged for 5 minutes at 4°C. The protein concentration in the supernatant was determined using a BCA Protein Assay Kit (Pierce, Rockford, IL, USA). Ninety-six-well multiter ELISA plates were coated in triplicate with 5 μg of protein sample/well [36]. The plates were incubated overnight at room temperature. After coating, the plates

were washed three times with 0.15 mol/L NaCl and 0.05% Tween 20 washing solution. After washing, the plates were blocked with 2% BSA and 0.1% Tween 20 in PBS buffer for 30 minutes at 37°C. After blocking, the plates were rewashed with washing solution and then incubated with rabbit anti-type I collagen (1:1000) in incubation buffer for 1 hour at room temperature. Preimmune serum was used as control. The plates were re-washed and incubated with alkali phosphatase-conjugated antirabbit IgG (1:1000) (Sigma Chemical Co., St. Louis, MO, USA) in incubation buffer for 1 hour at room temperature. After washing thoroughly, substrate, disodium p-nitrophenyl phosphate (5 µg/mL) (Sigma Chemical Co.) was added, and the absorbance was measured with an ELISA autoreader (Microplate Reader Model 3550) (Bio-Rad) at 450 nm.

Plasma biochemistry

At the end of the study, plasma samples collected from 3 mice in each group that did not receive PET were analyzed biochemically. The serum albumin levels were determined with the bromocresol green method (A/G B test, Wako, Osaka, Japan). Blood urea nitrogen, creatinine, total cholesterol, alanine aminotransferase, and alkaline phosphatase levels were measured with an auto-analyzer (Dry-Chem 3000, Fuji, Tokyo, Japan).

Modified PET

Modified PET was carried out to determine the ultrafiltration capacity of peritoneum following PFS. Five milliliters of 7% glucose dialysis fluid was instilled into the peritoneal cavity and maintained there for 2 hours. At the end of the PET, each mouse was sacrificed by cervical dislocation under ether anesthetic. Remaining CAPD fluid was drained through the catheter, and then laparotomy was performed to collect residual intraperitoneal fluid and tissue samples. The drainage volume was weighed, and the glucose concentration was determined with Dry-Chem 3000.

Statistical analysis

Values were presented as means \pm SE. Statistical differences between groups were evaluated by analysis of variance, followed by Duncan's multiple range test, with $P < 0.05$ used to detect significance.

RESULTS

General observations in mice with PFS

Chronic administration of saline for 4 weeks in control mice (Fig. 1) did not significantly alter the visceral peritoneum (representative section; Fig. 2A). Although chronic administration of CHG saline into the peritoneal cavity was not lethal, most PFS mice did not gain weight during the 4 weeks study period. After 2 weeks of every other

day administration of CHG, the visceral peritoneal histology from PFS mice demonstrated a mesothelium largely intact with a thickened subserosa. By week 4, the subserosal peritoneal tissue of PFS mice was significantly thickened with fibrocellular components (representative sections; Fig. 2 B and C); FSP1⁺ fibroblasts were beginning to accumulate at week 2 (Fig. 2E) and in greater numbers by week 4 (Fig. 2F) compared to the subserosa of control mice (Fig. 2D). Most of the other cells in the subserosa of PFS mice were Mac-3⁺ macrophages (Fig. 2G); Mac-3⁺ cells were FSP1⁻ by FACSScan (data not shown). CD4⁺ and CD8⁺ T lymphocytes were only identified occasionally in the subserosal peritoneum of PFS mice (data not shown). Furthermore, increased deposition of type I collagen and an increased number of CD31⁺ vascular endothelial cells in the subserosal peritoneum of PFS mice were found in fibrotic areas (Fig. 2 H and I). The numbers of FSP1⁺ fibroblasts and Mac-3⁺ macrophages, the amount of type I collagen and the extent of vascularity in the subserosal tissue each increased over time ($P \leq 0.05$) (Fig. 3). Although the number of Mac-3⁺ macrophages were fewer than the number of FSP1⁺ fibroblasts throughout the study, the macrophages invaded the subserosal tissue at weeks 1 to 2 before the fibroblasts increased in the number (Fig. 3 A and B). The presence of proliferating fibroblasts and infiltrating macrophages was followed by increasing deposits of type I collagen and neoangiogenesis (Fig. 3 C and D). Dual immunocytochemistry of the subserosal peritoneum of PFS mice revealed that most of the MCP-1, HSP47, or VEGF⁺ stromal cells were simultaneously positive for FSP1 (Fig. 4 A to C), suggesting that peritoneal fibroblasts play a role in the enrollment of macrophages, collagen production, and neoangiogenesis in PFS. Overall, the peritoneal histology in control mice was nearly normal, and chronic insertion of the catheter itself, at least for 4 weeks, did not cause any substantial changes to the visceral peritoneum. All the controls for immunohistochemistry were without significant background signals.

Deletion of fibroblasts in mice with PFS

In preliminary studies, we evaluated several general parameters prior to undertaking the deletion experiments. First, flow cytometric analysis of splenocytes derived from FSP1. Δ tk mice, with or without GCV treatment, revealed no remarkable alterations in the cellular composition of the spleen. For example, the percentage of CD4⁺ and CD8⁺ T lymphocytes, and Mac-3⁺ macrophages from FSP1. Δ tk mice receiving GCV were 27% \pm 2%, 14% \pm 2%, and 27% \pm 3%, respectively, and 23% \pm 2%, 11% \pm 2%, 24% \pm 2%, respectively, in FSP1. Δ tk mice not receiving GCV. Second, we observed in peritoneal fibrosis-transgenic mice that fibroblasts in the peritoneum at week 2 were simultaneously positive for Δ tk protein and FSP1 (Fig. 4d). Third, a nonlethal dose of

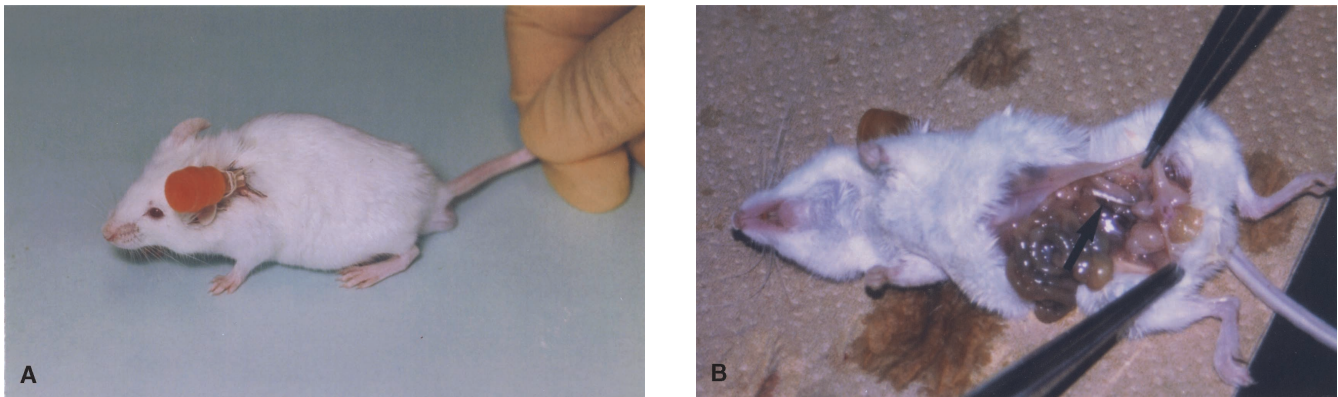


Fig. 1. Murine model of peritoneal fibrosing syndrome (PFS). (A) A 21 gauge plastic catheter was inserted from the back into the peritoneal cavity, fixed to the back skin, and maintained for 4 weeks. Each mouse was kept individually in one cage. (B) The catheter tip was located intraperitoneally (arrow). After each insertion procedure, normal saline was infused and drained through the catheter to confirm the location of the catheter tip.

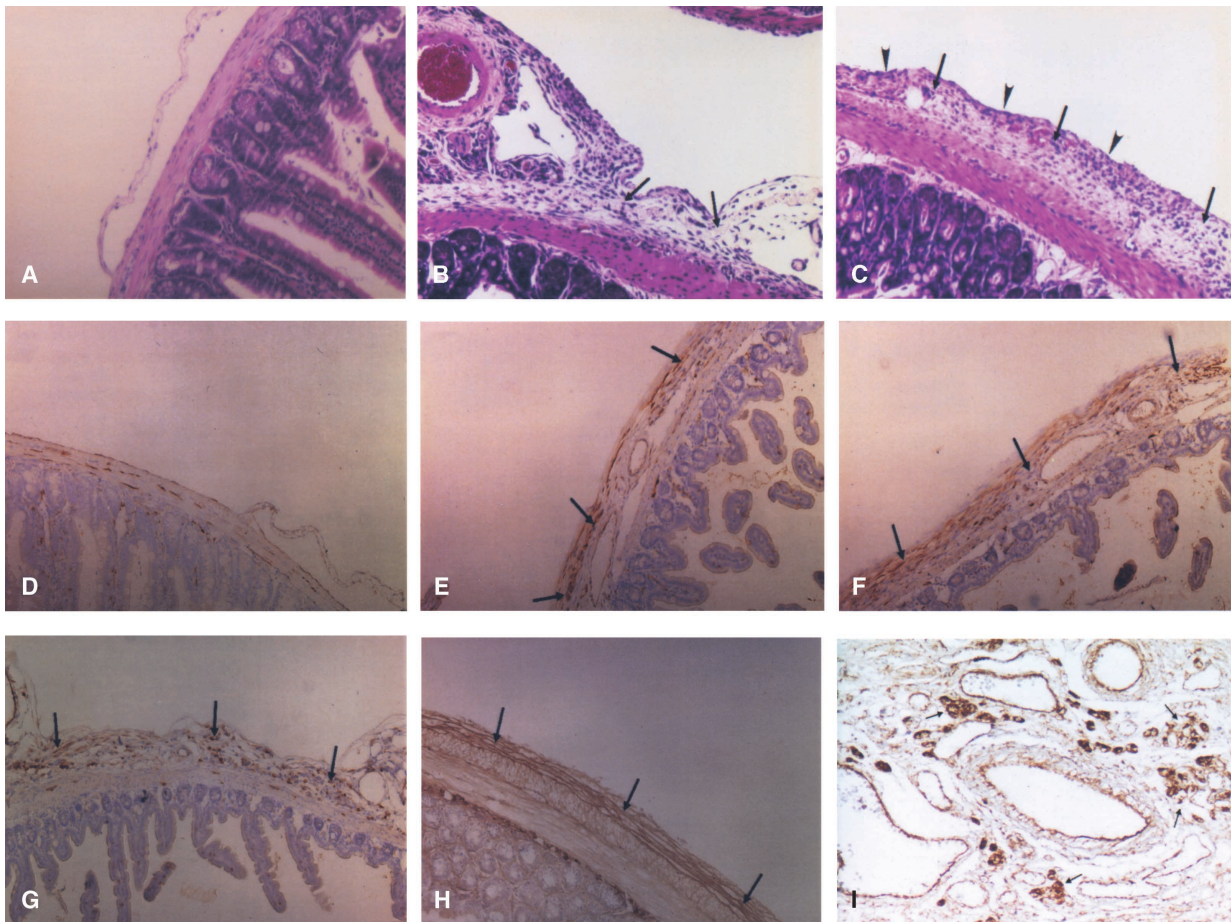


Fig. 2. Typical histopathology of murine peritoneal fibrosing syndrome (PFS). (A) Visceral peritoneum of control-wild-type mice at week 4. Subserosal stroma was thin and few cellular components were observed (hematoxylin-eosin stain, 150 \times). (B) Visceral peritoneum of peritoneal fibrosis-wild-type mice at week 2. Subserosal stroma was thickened (arrows) and contained cellular components (hematoxylin-eosin stain, 150 \times). (C) Visceral peritoneum of peritoneal fibrosis-wild-type mice at week 4. Subserosal stroma was significantly thickened (arrows) and both cellular and acellular areas were observed. Much of mesothelium remained intact (arrowheads) (hematoxylin-eosin stain, 150 \times). (D) There were few fibroblast-specific protein-1 (FSP1⁺) fibroblasts in the subserosal stroma of control-wild-type mice at week 4 diaminobenzidine (DAB) (150 \times). (E) There were numerous FSP1⁺ fibroblasts in the subserosal stroma (arrows) of peritoneal fibrosis-wild-type mice at week 2 (DAB, 150 \times). (F) These FSP1⁺ fibroblasts in the cellular area of the subserosal stroma (arrows) remained in peritoneal fibrosis-wild-type mice at week 4 (DAB, 150 \times). (G) There were a number of Mac-3⁺ macrophages in the subserosal stroma (arrows) of peritoneal fibrosis-wild-type mice at week 2 (DAB, 150 \times). (H) There was significant deposition of type I collagen in the subserosal stroma (arrows) in peritoneal fibrosis-wild-type mice at week 4 (DAB, 150 \times). (I) The thickened subserosal tissue in peritoneal fibrosis-wild-type mice at week 4 was rich in CD31⁺ small vessels (arrows) (DAB, 200 \times).

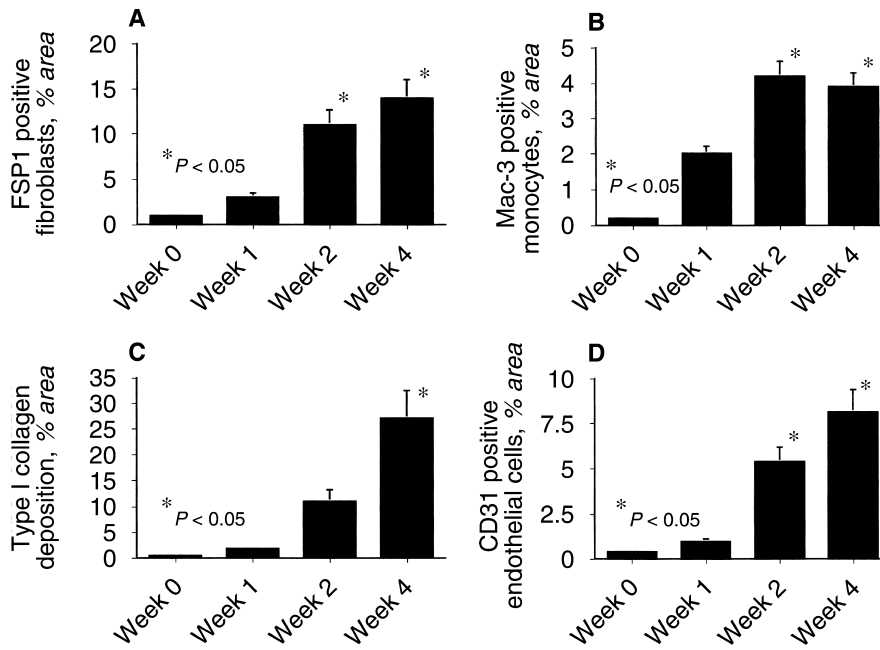


Fig. 3. Quantitative assessment of fibroblast-specific protein-1 (FSP1⁺) fibroblasts, Mac-3⁺ macrophages, and extent of type I collagen deposition in peritoneal fibrosis-wild-type mice. (A) The number of FSP1⁺ fibroblasts in the subserosal stroma significantly increased over time. (B) The number of Mac-3⁺ macrophages in the subserosal stroma significantly increased by week 2 and remained until week 4. (C) The amount of type I collagen deposition in the subserosal stroma of the peritoneum increased significantly by week 4. (D) The hypervascularity in the subserosal stroma was significant at week 4.

GCV (50 mg/kg) once a day by catheter infusion was well tolerated for a treatment protocol of 2 weeks (data not shown). Finally, the metabolic profile of mice from all groups was not significantly different from each other (data not shown).

In experiments to formally test the role of peritoneal fibroblasts in mice with PFS, the visceral peritoneum of peritoneal fibrosis-wild-type + GCV mice after 4 weeks of CHG saline were fibrotic and significantly thickened compared to peritoneal fibrosis-transgenic + GCV mice (Fig. 5 A and B), and seemed substantially similar to peritoneal fibrosis mice in the pilot study. Apoptosis was induced exclusively in FSP1⁺ fibroblasts (Fig. 4E) in peritoneal fibrosis-transgenic mice with exposure to GCV such that the number of FSP1⁺ fibroblasts in the peritoneum was significantly reduced below those in peritoneal fibrosis-wild-type mice. The percentage of the apoptotic nuclei was $45.5\% \pm 5.5\%$ among 40 ± 4 FSP1⁺ fibroblasts observed in a $300\times$ field of subserosal stroma from peritoneal fibrosis-transgenic mice at week 4, but only $2.0\% \pm 0.4\%$ among 49 ± 5 FSP1⁺ fibroblasts in peritoneal fibrosis-wild-type mice ($P < 0.05$) (Figs. 5 C and D and 6 A).

When the proliferation of FSP1⁺ fibroblasts was attenuated in peritoneal fibrosis-transgenic mice receiving GCV, the number of Mac-3⁺ macrophages and CD31⁺ endothelial cells, and the accumulation of type I collagen were also reduced in the peritoneum of peritoneal fibrosis-transgenic mice ($P \leq 0.05$) (Figs. 5 E to I and 6 B to D), leaving the subserosal tissues of the visceral peritoneum in peritoneal fibrosis-transgenic mice considerably intact. The peritoneum of control-transgenic mice and control-wild-type mice had no significant abnormalities (data not

shown). Quantification analyses by indirect ELISA also demonstrated that deletion of fibroblasts in peritoneal fibrosis-transgenic mice treated with GCV reduced type I collagen deposition to control levels ($P \leq 0.05$) (Fig. 7).

Modified PET

At the conclusion of the 4 week study, each group of mice (peritoneal fibrosis-transgenic, peritoneal fibrosis-wild-type, control-transgenic, and control-wild-type) underwent a modified PET. The drainage volume and glucose concentration of each group were measured (Fig. 8). These indices were significantly lower in peritoneal fibrosis-wild-type + GCV mice compared to those measured in peritoneal fibrosis-transgenic + GCV mice ($P \leq 0.05$), suggesting that selective deletion of fibroblasts in peritoneal fibrosis-transgenic mice reduced glucose absorption and preserved ultrafiltration.

DISCUSSION

We observed that the number of FSP1⁺ fibroblasts in the subserosal stroma of mice with PFS increased significantly over a 4-week interval following chronic exposure to CHG saline. At week 2, the PFS lesions generated in peritoneal fibrosis-transgenic mice showed increasing numbers of FSP1⁺ fibroblasts. GCV treatment started at that time for an additional 2 weeks rescued the peritoneum from PFS by decreasing the number of fibroblasts to the control levels via apoptosis. Our approach to peritoneal rescue was associated with a decrease in monocyte infiltration, collagen deposition, and neoangiogenesis.

Since studies of fibroblasts in vivo have been limited

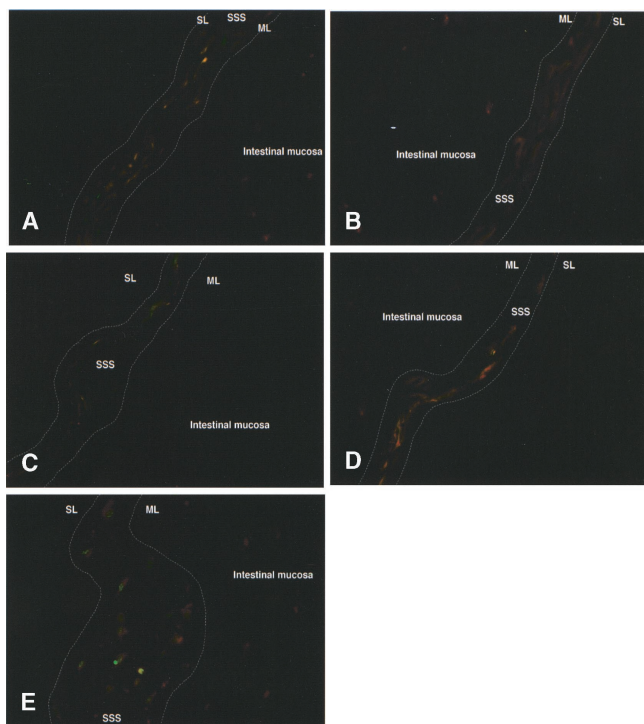


Fig. 4. Characterization of peritoneal fibroblast-specific protein-1 (FSP1⁺) fibroblasts in the subserosal stroma. (A) Immunofluorescence of FSP1 (in red) and monocyte chemoattractant protein-1 (MCP-1) (in green) in peritoneal fibrosis-wild-type mice at week 4. There were some stromal cells in yellow in the subserosal stroma, indicating that some fibroblasts produced MCP-1 (merged image, 150 \times). (B) Using two labels, FSP1 (in red) and heat shock protein 47 (HSP47) (in green), there were some stromal cells in yellow in the subserosal stroma of peritoneal fibrosis-wild-type mice at week 4, indicating that FSP1⁺ fibroblasts contributed to collagen production (merged image, 150 \times). (C) Immunofluorescence of FSP1 (in red) and vascular endothelial growth factor (VEGF) (in green) in peritoneal fibrosis-transgenic mice at week 2. Some of FSP1⁺ fibroblasts in the subserosal stroma (in yellow) were simultaneously positive for VEGF, suggesting their contribution to neoangiogenesis (merged image, 150 \times). (D) Immunofluorescence of FSP1 (in red) and thymidine kinase (tk) protein (in green) in peritoneal fibrosis-transgenic mice at week 2. FSP1⁺ fibroblasts in the subserosal stroma were simultaneously positive for tk protein, yielding yellow color (merged image, 150 \times). (E) Many of the FSP1⁺ fibroblasts were selectively killed by gancyclovir (GCV) treatment via apoptosis (arrows) on colocalization of FSP1 expression (in red) and apoptotic nuclei (in green) in peritoneal fibrosis-transgenic mice at week 4 following GCV treatment (merged image, 300 \times). Abbreviations are: SL, serosal layer; SSS, subserosal stroma; ML, muscular layer.

by a lack of fibroblasts-specific markers, we took advantage of the marker, FSP1 [27, 29]. FSP1 belongs to the S-100 superfamily and is associated with the regulation of cell shape and motility [27, 36]. Using anti-FSP1 antibody, we have previously demonstrated that the number of FSP1⁺ fibroblasts increase in parallel with the progression of renal fibrosis [27, 28, 31]. The same was true in the present study. Our FSP1. Δ tk transgenic mouse on exposure to GCV had some additional advantages, one of which is that GCV administration induces apoptosis

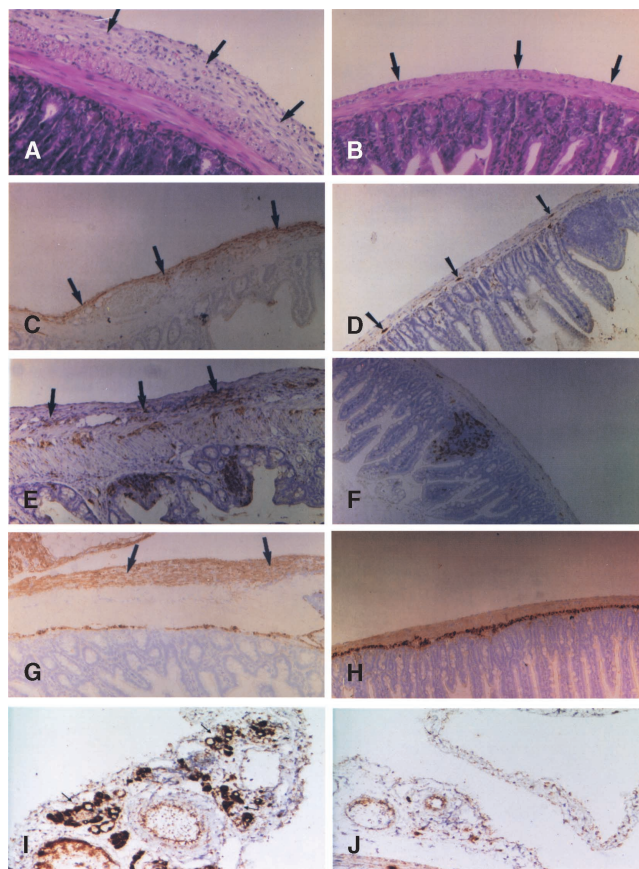


Fig. 5. Typical histopathology of peritoneal fibroblast-specific protein (PFS) in peritoneal fibrosis-transgenic and peritoneal fibrosis-wild-type mice treated with gancyclovir (GCV). (A) The subserosal stroma of the visceral peritoneum (arrows) in peritoneal fibrosis-wild-type mice was thickened and fibrous (hematoxylin-eosin stain, 150 \times). (B) With GCV treatment, the subserosal stroma of the visceral peritoneum (arrows) in peritoneal fibrosis-transgenic mice remained thin and contained few cellular components (hematoxylin-eosin stain, 150 \times). (C) There were a number of fibroblast-specific protein-1 (FSP1⁺) fibroblasts in the subserosal stroma (arrows) from peritoneal fibrosis-wild-type mice diaminobenzidine (DAB) (150 \times). (D) GCV treatment reduced the number of FSP1⁺ fibroblasts in the subserosal stroma (arrows) of peritoneal fibrosis-transgenic mice (DAB, 150 \times). (E) There were a number of Mac-3⁺ macrophages in the subserosal stroma (arrows) of peritoneal fibrosis-wild-type mice (DAB, 150 \times). (F) GCV treatment reduced the number of Mac-3⁺ macrophages in the subserosal stroma of peritoneal fibrosis-transgenic mice (DAB, 150 \times). (G) There was a significant deposition of type I collagen in the subserosal stroma (arrows) of peritoneal fibrosis-wild-type mice (DAB, 150 \times). (H) There was no significant deposition of type I collagen in the subserosal stroma of peritoneal fibrosis-transgenic mice (DAB, 150 \times). (I) Increased vascularity by CD31 staining was evident in the subserosal tissue of peritoneal fibrosis-wild-type mice (arrows) (DAB, 200 \times). (J) There was no significant vascularity in the subserosal tissue of peritoneal fibrosis-transgenic mice (DAB, 200 \times).

only in proliferating fibroblasts [31]. This approach works in connective tissues (peritoneum) as well as in complex organs [26, 31].

In the early days of CAPD in humans, it was believed that regular exposure of peritoneal membranes to anti-septics might reduce the risk of peritonitis [14]. However,

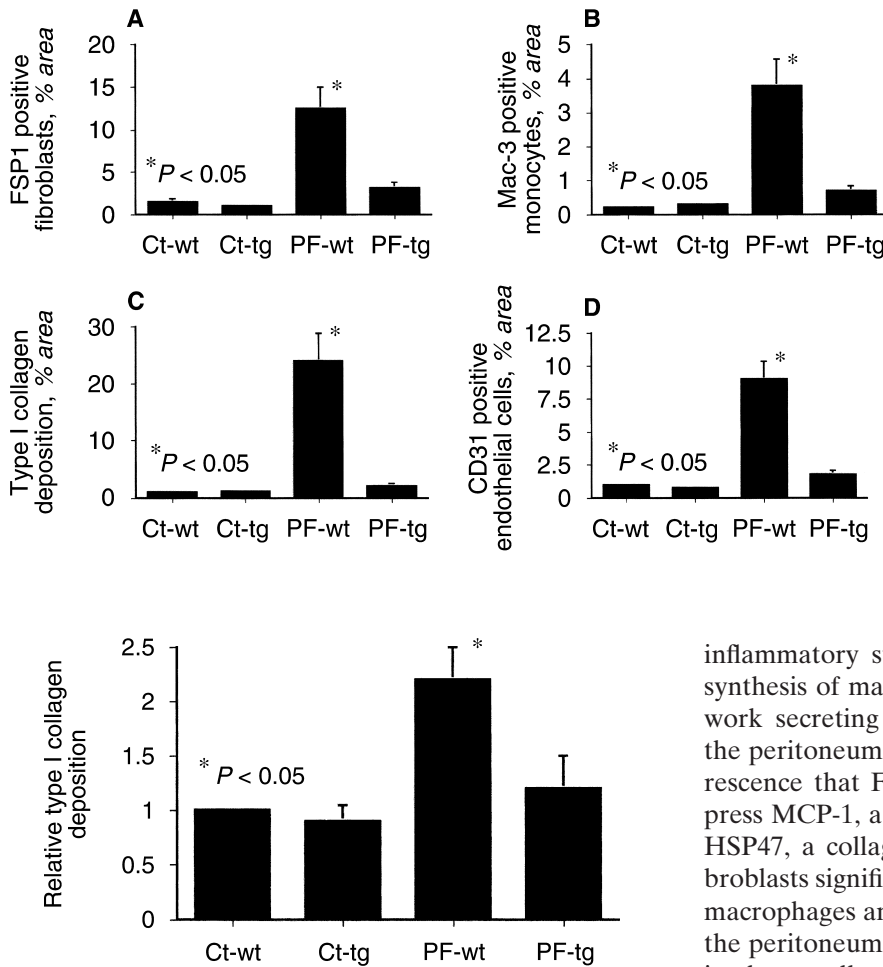


Fig. 6. Quantitative assessment of peritoneal fibroblast-specific protein (PFS) histopathology in peritoneal fibrosis-transgenic (PF-tg) and peritoneal fibrosis-wild-type (PF-wt) mice treated with gancyclovir (GCV). (A) The number of fibroblast-specific protein-1 (FSP1⁺) fibroblasts in peritoneal fibrosis-transgenic mice was significantly reduced to the control levels (Ct-wt and Ct-tg) by GCV treatment. (B) The number of Mac-3⁺ macrophages in peritoneal fibrosis-transgenic mice was reduced to control levels with GCV. (C) The amount of type I collagen deposited in the peritoneum of peritoneal fibrosis-transgenic mice was also maintained at control levels by deletion of fibroblasts. (D) Neovascularization was also inhibited in the subserosal tissue of peritoneal fibrosis-transgenic mice.

Fig. 7. Quantitative analysis of type I collagen in the peritoneal subserosa by indirect enzyme-linked immunosorbent assay (ELISA). The amount of type I collagen deposited in small bowel subserosa of peritoneal fibrosis-wild-type (PF-wt) mice treated with gancyclovir (GCV) was significantly greater than in peritoneal fibrosis-transgenic mice (PF-tg). Collagen in these peritoneal fibrosis-transgenic mice was significantly reduced to control levels (Ct-wt and Ct-tg) by deletion of fibroblasts following GCV treatment.

exposure to chlorhexidine in alcohol was identified early as responsible for an outbreak of sclerosing peritonitis [30]. This observation was confirmed by toxicologic experiments in animals [17, 19, 30, 37]. Accordingly, we introduced chlorhexidine into the peritoneum of mice to generate PFS [19, 30, 38]. Constant exposure of the mesothelium to chlorhexidine is thought to disrupt mesothelial tight junctions followed by crystalline damage to the subserosal stroma [14, 30]. Persistent serositis results from prolonged exposure to this or other toxic metabolites and creates an inflammatory response that predisposes to PFS.

Recent studies in CAPD patients suggest that peritoneal fibroblasts may derive from mesothelium by epithelial-mesenchymal transition [13]. Based on in vitro observations, peritoneal fibroblasts both respond to pro-

inflammatory stimuli with proliferation and increased synthesis of matrix, and contribute to the cytokine network secreting immunologically active molecules into the peritoneum [39]. We demonstrated by immunofluorescence that FSP1⁺ fibroblasts in the peritoneum express MCP-1, a macrophages chemoattractant [40], and HSP47, a collagen chaperone [41]. The deletion of fibroblasts significantly reduced the number of infiltrating macrophages and the accumulation of type I collagen in the peritoneum, suggesting that fibroblasts are involved in the enrollment of circulating macrophages and the production of type I collagen. At an early phase of PFS, macrophages are attracted by activated resident cells in the peritoneum, including mesothelial cells, fibroblasts, and infiltrated macrophages themselves [42–44]. Furthermore, fibroblasts likely anchor infiltrating macrophages, and fibroblast deletion subsequently erased the macrophages that would have otherwise infiltrated the subserosal tissue (Figs. 2G and 5F). These infiltrating macrophages can accelerate fibrogenesis by releasing profibrogenic cytokines and the fibroblasts growth factors described above [29, 45]. The reduction in their numbers might indirectly attenuate PFS. Our findings suggest that macrophages and fibroblasts cofacilitate each other, leading to a vicious cycle perpetuating fibrogenesis, and that deletion of FSP1⁺ fibroblasts can interrupt this cycle. The degree of experimental fibrosis may depend on the strain variability of mice. Compared to other strains, Balb/c mice (like we used in this study) may not form extensive fibrotic tissue under some experimental conditions [46–48]. We nevertheless observed significant peritoneal fibrosis in Balb/c mice.

We also observed enhanced glucose absorption and loss of ultrafiltration capacity in peritoneal fibrosis-wild-type mice treated with GCV. This functional deteriora-

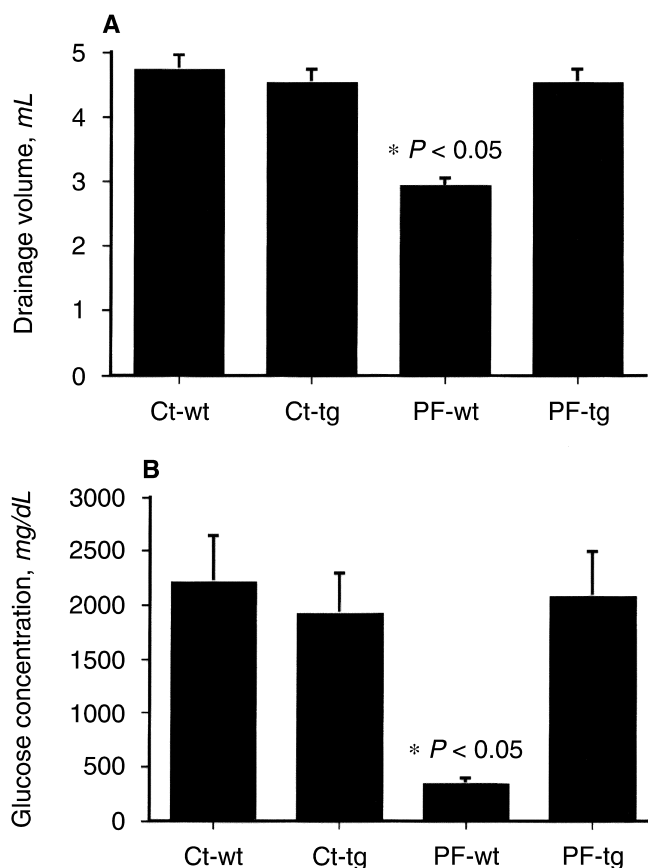


Fig. 8. Modified peritoneal equilibrium test (PET). (A) The volume of fluid drained from peritoneal fibrosing syndrome (PFS) mice. The volume of fluid drained from peritoneal fibrosis-wild-type (PF-wt) mice was significantly less than that of peritoneal fibrosis-transgenic mice (PF-tg) treated with gancyclovir (GCV) ($P \leq 0.05$), and similar to the control levels (Ct-wt and Ct-tg) in mice not treated with chlorhexidine gluconate (CHG) saline. This suggests that ultrafiltration capacity was preserved in peritoneal fibrosis-transgenic mice by deletion of fibroblasts compared to peritoneal fibrosis-wild-type mice. (B) The glucose concentration of drained fluid in peritoneal fibrosis-wild-type mice was significantly lower ($P \leq 0.05$) than that of peritoneal fibrosis-transgenic mice treated with GCV, and similar to the control levels in mice not treated with CHG saline. This suggests that the ultrafiltration failure observed in peritoneal fibrosis-wild-type mice was due to the increased diffusion of glucose through the peritoneal membrane.

tion of the peritoneum is similar to the same well-known complication of long-term CAPD. Several different mechanisms for loss of filtration across the peritoneum have been described [1, 49–51]. Some authors emphasize the role of increased solute transport to explain the loss of ultrafiltration due to peritoneal hypervascularity [52, 53], whereas others suggest the possible impact of changing lymphatic or peritoneal fluid absorption into peritoneal tissue [54, 55]. A third possibility is that the formation of adhesions and sclerosis results in a hyperpermeable peritoneum even with loss of surface area [1]. Finally, impaired efficiency of the osmotic agent in peritoneal dialysate, without change in solute transport, may contribute to loss of ultrafiltration, perhaps related to alter-

ations in transcellular water channels (aquaporin-1) in the vascular endothelium [56–58]. Among these mechanisms, enhanced glucose absorption due to hypervascularity in the peritoneal membrane is the most common explanation for loss of ultrafiltration in CAPD [51–53].

In the present study, we also observed that fibroblasts express VEGF, an angiogenic growth factor, which might be involved in neoangiogenesis of subserosal tissue [53, 59]. Several investigators reported that fibroblasts producing VEGF provide a basic framework for proliferating endothelial cells that form vascular structures [60–62]. Therefore, antifibrotic therapy by deleting FSP1⁺ fibroblasts may be important in the prevention of CD31⁺ neoangiogenesis in subserosal tissue. We suggest that controlling the number of FSP1⁺ fibroblasts assists in maintaining mesothelial architecture, while reducing the infiltration of macrophages, collagen accumulation, and neoangiogenesis that preserves peritoneal function.

Although CAPD patients are not given chlorhexidine any more, there are a growing number of patients on CAPD suffering from PFS and ultrafiltration failure [1–5, 12, 14]. The histopathologic findings in our model of PFS are similar to the peritoneal characteristics of CAPD patients after a couple of episodes of peritonitis [12, 14, 63] and supports the clinical relevance of our experimental system. A few experimental trials to prevent PFS using pirfenidone or glycosaminoglycans have been reported [11, 19]. To our knowledge, it is the first study demonstrating that peritoneal fibroblasts play a pivotal role in the process of PFS. Their conditional deletion has therapeutic potential.

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